

CLAIMS

This listing of the claims will replace all prior versions, and listings, of the claims in the application:

Listing of Claims

1-23. (Cancelled)

24. (Currently Amended) A method of treatment of or protection against an inflammatory disease comprising administering an effective amount of a peptide of 7-30 amino acids derived from the sequence 241-270 of SEQ ID NO. 1 representing the sequence of the stress protein hsp65 of *Mycobacterium tuberculosis*, wherein at least a portion of the peptide comprises a sequence selected from the group consisting of PLXIIAE, LXIIAED, XIIAEDV, IIAEDVX, IAEDVXG, AEDVXGE, DVXGEAL, VXGEALS, XGEALST, GEALSTL, EALSTLV, ALSTLVVALSTLVX, LSTLVXN and STLVXNX, and wherein X is any amino acid.

25-27. (Cancelled)

28. (Currently Amended) The method of claim 24, wherein the peptide comprises the amino acids derived from the sequence 256-265 of SEQ ID NO. 1, wherein the portion of the peptide has the sequence selected from the group consisting of EALSTLV, ALSTLVVALSTLVX, LSTLVXN and STLVXNX.

29. (Previously Presented) The method of claim 24, wherein one or more of the amino acids residues has been exchanged with a residue of an amino acid having similar size, charge and polarity.

30. (Cancelled)

31. (Previously Presented) The method of claim 24, wherein the peptide is administered parenterally, orally or nasally.

32. (Previously Presented) The method of claim 31, wherein the peptide is administered nasally.

33. (Previously Presented) The method of claim 25, wherein the peptide is administered parenterally, orally or nasally.

34. (Currently Amended) The method of claim 28, wherein the portion of the peptide has the sequence selected from the group consisting of EALSTLV, ALSTLVVALSTLVX, LSTLVVN and STLVVNK.

35. (Previously Presented) The method of claim 28, wherein the portion of the peptide has the sequence selected from the group consisting of EALSTLV, ALSTLVVALSTLVX, LSTLVLN and STLVVLR.

36. (New) The method of claim 24, further comprising administering an effective amount of a second peptide of 7-30 amino acids derived from the sequence 241-270 of SEQ ID NO. 1 representing the sequence of the stress protein hsp65 of *Mycobacterium tuberculosis*, wherein at least a portion of the peptide comprises a sequence selected from the group consisting of PLXIIAE, LXIIAED, XIIAEDV, IIAEDVX, IAEDVXG, AEDVXGE, DVXGEAL, VXGEALS, XGEALST, GEALSTL, EALSTLV, ALSTLVX, LSTLVXN and STLVXNX, wherein X is any amino acid, and wherein the second peptide is different from the peptide.